

# Nuclear Receptors License Phagocytosis by Trem2<sup>+</sup> Myeloid Cells in Mouse Models of Alzheimer's Disease

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Alzheimer's disease (AD) is characterized by a robust inflammatory response elicited by the accumulation and subsequent deposition of amyloid (A $\beta$ ) within the brain. The brain's immune cells migrate to and invest their processes within A $\beta$  plaques but are unable to efficiently phagocytose and clear plaques from the brain. Previous studies have shown that treatment of myeloid cells with nuclear receptor agonists increases expression of phagocytosis-related genes. In this study, we elucidate a novel mechanism by which nuclear receptors act to enhance phagocytosis in the AD brain. Treatment of murine models of AD with agonists of the nuclear receptors PPAR $\gamma$ , PPAR $\delta$ , LXR, and RXR stimulated microglial phagocytosis *in vitro* and rapidly induced the expression of the phagocytic receptors Axl and MerTK. In murine models of AD, we found that plaque-associated macrophages expressed Axl and MerTK and treatment of the cells with an RXR agonist further induced their expression, coincident with the rapid reduction in plaque burden. Further characterization of MerTK<sup>+</sup>/Axl<sup>+</sup> macrophages revealed that they also expressed the phagocytic receptor TREM2 and high levels of CD45, consistent with a peripheral origin of these cells. Importantly, in an *ex vivo* slice assay, nuclear receptor agonist treatment reversed the AD-related suppression of phagocytosis through a MerTK-dependent mechanism. Thus, nuclear receptor agonists increase MerTK and Axl expression on plaque-associated immune cells, consequently licensing their phagocytic activity and promoting plaque clearance.

**Key words:** Alzheimer's disease; microglia; nuclear receptors; phagocytosis

## Introduction

Inflammation plays critical roles in the initiation and progression of Alzheimer's disease (AD) (Wyss-Coray, 2006). This has been highlighted by genetic studies that identify numerous immune mediators linked to AD risk (Jones et al., 2010; Hollingworth et al., 2011; Naj et al., 2011). Notably, variants of the microglial cell surface receptor TREM2 confer dramatically elevated risk for AD and other neurodegenerative diseases (Cuyvers et al., 2014). The AD brain is typified by the presence of abundant "activated" inflammatory macrophages that are found principally associated

with extracellular deposits of amyloid (Akiyama et al., 2000). An enigmatic feature of the disease is that these "activated" macrophages are unable to mount a phagocytic response directed toward amyloid deposits, despite expression of phagocytic receptors, including TREM2, Axl, and MerTK. This results in the progressive accumulation of plaques within the brain.

The phenotypes of microglia and other myeloid cells are governed by the actions of heterodimeric Type II nuclear receptors, most prominently PPAR $\gamma$ :RXR, PPAR $\delta$ :RXR, and LXR:RXR (Glass and Saijo, 2010). These ligand-activated transcription factors act to "license" complex cellular responses through their ability to coordinately regulate arrays of functionally allied genes (Hucke et al., 2012, 2013). In myeloid cells, nuclear receptors drive the acquisition of "alternative activation" states that are responsible for the resolution of inflammation (Glass et al., 2010), and the promotion of tissue repair and phagocytosis (Nagy et al., 2012). In murine models of AD, nuclear receptor agonist treatment suppresses inflammation, reduces amyloid accumulation, and improves cognition (Skerrett et al., 2014). The reduction in plaque burden is due to nuclear receptor stimulation of phagocytosis, and the present study implicates the phagocytic receptors Axl and MerTK in this response. Importantly, the expression of MerTK, Axl (Mukundan et al., 2009; A-Gonzalez et al., 2009), and TREM2 (Daniel et al., 2014) is directly regulated by nuclear receptor agonists.

The phagocytic receptors TREM2, MerTK, and Axl are expressed on myeloid cells, including microglia (Gautier et al.,

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2012; Butovsky et al., 2014). Axl and MerTK are TAM-family receptor tyrosine kinases that are required for the phagocytosis of apoptotic cells (Lemke, 2013). These receptors recognize exposed phosphatidylserine on the surface of apoptotic cells through the adaptor molecules Protein S or Gas6 (Hafizi and Dahlbäck, 2006). In tissue, Gas6 is often constitutively bound to Axl but does not signal in the absence of phosphatidyl serine (Zagórska et al., 2014). Gas6 binds to and can activate all TAM receptors, whereas Protein S cannot activate Axl efficiently, and the relevant ligand domains have recently been elucidated (Lew et al., 2014). TREM2 has been postulated to be a phagocytic receptor whose expression is also associated with suppression of inflammatory gene expression (Turnbull et al., 2006; Hsieh et al., 2009; N'Diaye et al., 2009).

Although MerTK, Axl, and TREM2 are expressed on plaque-associated macrophages in murine models of AD, these cells are phagocytically inactive. MerTK- and TREM2-expressing macrophages are also CD45<sup>hi</sup>, a canonical marker of peripherally derived “inflammatory” monocytes (Sedgwick et al., 1991). In allied work (Jay et al., 2015), the presence of plaque-associated macrophages in the AD brain was reliant upon the expression of TREM2, which was found to be exclusively expressed by CD45<sup>hi</sup> macrophages. Importantly, nuclear receptor activation enables phagocytic machinery within brain macrophages, licensing phagocytosis, resulting in the reduction in plaque burden *in vivo*.

## Materials and Methods

**Animals.** B6C3-Tg (APP/PS1Δe9) 85Dbo/J mice (Jankowsky et al., 2001) as well as B6SJL-Tg(5XFAD) mice (Oakley et al., 2006) were obtained from The Jackson Laboratory. At 12 months of age, APP/PS1Δe9 animals were treated with 100 mg/kg bexarotene (Targretin) or water by oral gavage for 5 d before slice phagocytosis, or 7 d (Cramer et al., 2012) before death. Targretin (75 mg) was dispersed in H<sub>2</sub>O to a concentration of 30 mg/ml, and animals received 100 μl of solution by oral gavage, resulting in a treatment of 100 mg/kg/d. At 4 months of age, 5XFAD mice were treated with 100 mg/kg/d bexarotene, 30 mg/kg/d GW0742 (GlaxoSmithKline), or water for 14 d. At 8 months of age, 5XFAD mice were treated with 100 mg/kg/d bexarotene for 7 d. An equal number of male and female animals were used for each experiment. Animals were housed and cared for according to Case Western Reserve University's Institutional Animal Care and Use Committee committee.

**Nuclear receptor agonists.** The following nuclear receptor agonists were used: LXR agonist GW3965, PPAR $\gamma$  agonist pioglitazone, PPAR $\delta$  agonist GW0742, and RXR agonist bexarotene. Drugs were dispersed in water for oral gavage (GW0742, bexarotene) or dissolved in DMSO for tissue culture.

**Tissue collection.** Four hours after their last gavage, animals were killed and transcardially perfused with saline. The left hemispheres were fixed overnight in 4% PFA in phosphate buffer, while the cortices and hippocampi were dissected from the right hemispheres and processed for protein and A $\beta$  extraction. The tissue was homogenized in tissue homogenization buffer (20 mM Tris, pH 7.4, 250 mM sucrose, 0.5 mM EDTA, 0.5 mM EGTA) using a tissue homogenizer. A sequential extraction of A $\beta$  was performed and A $\beta$ 40 and 42 levels were analyzed by ELISA as previously described (Jiang et al., 2008). A second cohort of 12-month-old animals were processed for adult microglial isolation.

**Microglial/macrophage isolation, RNA extraction, and qRT-PCR.** Adult microglia/macrophages were isolated as described previously (Cardona et al., 2006). Briefly, cortices and hippocampi were removed and dissociated using a sterile razorblade and enzymatically digested in RPMI media containing collagenase Type 3 (0.2%, Worthington) and dispase (2 U/ml, Invitrogen) for 45 min at 37°C. Digestion was stopped by adding 15 ml of PBS containing 2.5 mM EDTA and 1% FBS. Homogenates were passed through a 35 μm filter and centrifuged. Pellets were resuspended in a 30% Percoll solution and centrifuged to separate microglia and macrophages from other cell types. The purity of CD11b<sup>+</sup>/CD45<sup>+</sup> mac-

rophages was ~70% as assessed by flow cytometry. After several PBS washes, cells were pelleted and frozen immediately at -80°C for mRNA extraction. mRNA was extracted using RNeasy spin columns (QIAGEN), genomic DNA was eliminated, and mRNA was reverse transcribed using a quantitect reverse transcription kit (QIAGEN). The resulting cDNA was preamplified using TaqMan preamp master mix (Applied Biosystems) and detected using a Step One plus and TaqMan assays (Applied Biosystems). Data presented represent relative quotient with 95% confidence intervals, and statistical analyses were performed on  $\Delta$ Ct values (Yuan et al., 2006).

**Flow cytometry.** Adult microglia and macrophages were isolated and resuspended in FACS buffer (1% FBS in PBS) containing FC block for 15 min before incubation with appropriate antibodies. Cells were incubated with tagged antibodies for 30 min before washing, fixing, and analysis using a FACSAria. Analysis was done using FlowJo software. Cells were gated on CD11b<sup>+</sup> populations and further separated into CD45<sup>lo</sup> and CD45<sup>hi</sup> cells. The following antibodies were used: anti-mouse TREM2 (R&D Systems, clone 237920), anti-mouse MerTK (R&D Systems, clone 108928), anti-mouse CD45 (eBioscience, clone 30-F11), and anti-mouse CD11b (eBioscience, clone M1/70).

**Immunohistochemistry.** Tissue sections (10 μm) were rehydrated in PBS, followed by antigen retrieval in 10 mM sodium citrate or Reveal Decloaker (Biocare Medical) at 80°C for 20 min. Sodium citrate was used for all images where fluorescence intensity was measured, whereas Reveal Decloaker was used for TREM2 and confocal images. Sections were incubated in blocking solution (2% donkey serum and 0.1% Triton X-100 in PBS) for 1 h followed by incubation with primary antibody in blocking solution overnight at 4°C. Following several PBS rinses, sections were incubated with appropriate secondary antibodies (Invitrogen) and stained with DAPI. TREM2 fluorescence was visualized using tyramide amplification according to the manufacturer's instructions (PerkinElmer). Images were analyzed using ImageJ software. Mean fluorescence intensity of plaque-associated cells was measured and normalized to the vehicle-treated animals. The experimenter was blinded to genotype and treatment groups. The following antibodies were used: anti-mouse TREM2 (LSBio), anti-mouse MerTK (R&D Systems), anti-mouse Axl (R&D Systems), anti-mouse Gas6 (R&D Systems), anti-mouse Iba1 (Wako), and anti-mouse CD45 (BD Biosciences).

**Cell culture.** The N9 mouse microglial cell line was grown in RPMI containing 5% heat-inactivated FBS and 1% penicillin/streptomycin. Cells were plated at a density of 500,000 cells/well in a 6-well plate for *in vitro* phagocytosis assays. Primary microglia were cultured from P0–3 C57BL/6j pups as previously described (McDonald et al., 1997). Briefly, neonatal pups were killed, and their brains removed, mechanically dissociated, and digested in 0.5% Trypsin-EDTA for 20 min at 37°C. Digestion was stopped by addition of 10% FBS in DMEM/F12 media containing 1% penicillin/streptomycin, and the homogenate was plated on 150 mm plates and allowed to grow for 14–21 d at 37°C and 5% CO<sub>2</sub>. Microglia were separated using a mild trypsin protocol (Saura et al., 2003) and switched to serum-free media 24 h before drug treatments.

**Western blotting.** Protein levels of cell lysates or brain extracts were measured by BCA, and equal amounts of protein from each sample were resolved on 4%–12% Bis/Tris gels (Invitrogen). The following primary antibodies were used: goat anti-MerTK (R&D Systems), goat anti-Mouse Gas6 (R&D Systems), and goat anti- $\beta$ -actin (Santa Cruz Biotechnology).

**In vitro phagocytosis assay.** Amyloid was fibrillized by dissolving 1 mg A $\beta$ <sub>1–42</sub> in 220 μl endotoxin-free water and incubating for 5 d at 37°C. N9 or primary microglia were incubated with nuclear receptor agonists in serum-free media for 18 h. Following drug pretreatment, cells were treated  $\pm$ 2.5 μM fA $\beta$  or 40 nM Gas6 for 15 min before incubation with 1 μM Nile red-labeled polystyrene beads (Invitrogen) for 15 min. Immune IgG was used as a positive control to drive phagocytosis (Koenigsnecht-Talboo and Landreth, 2005). Cells were washed thoroughly with PBS, fixed in 2% PFA in PBS, and phagocytic cells were measured via flow cytometry using a BD FACSAria. Data were analyzed using FlowJo software and normalized to nontreated control cells.

**Ex vivo phagocytosis assay.** Phagocytosis assays were performed in brain slices as described previously (Krabbbe et al., 2013). Briefly, mice were treated with vehicle or 100 mg/kg/d bexarotene for 5 d before death.

Mice were lightly anesthetized with isoflurane, killed by cervical dislocation, and brains were removed and 150  $\mu\text{m}$  slices prepared using a Leica vibratome. After resting in RPMI for 30–60 min, slices were incubated with 2  $\mu\text{m}$  fluorescent beads alone or in the presence of recombinant Gas6 or MerTK function blocking antibodies. Following incubation, slices were washed for 1 h, fixed, and stained using anti-Iba1 to identify myeloid cells. Slices were imaged on a Leica confocal microscope and analyzed using ImageJ. The number of beads per cell were counted, and the phagocytic index was calculated by determining the percentage of cells containing 0, 1–4, 5–7, 8–10, or >10 beads. The percentage of cells in each group was then multiplied by the grade of phagocytosis (1–4:1, 6–7:2, 8–10:3, >10:4), and the sum of the products in each group was then normalized to values obtained from the nontransgenic animal. The experimenter was blinded to mouse genotype and treatment.

**Statistical analysis.** All values reported as mean  $\pm$  SEM. Statistics were evaluated using Graphpad Prism software and Student's *t* test or one-way ANOVA with Tukey's *post hoc* where noted.

## Results

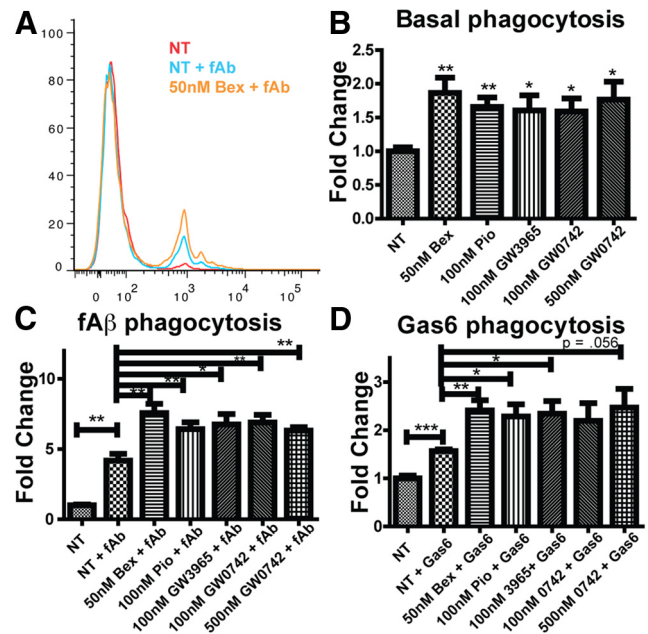
We use the term “microglia” to refer to resident myeloid cells of the CNS parenchyma that are derived from yolk sac progenitors that invade the neuroepithelium early (E8.5) in development (Ginhoux et al., 2010; Schulz et al., 2012), whereas “macrophages” refer to activated parenchymal myeloid cells, which can be derived either from peripherally derived CD45<sup>hi</sup> monocytes or resident microglial cells (Ransohoff and Cardona, 2010; Mildner et al., 2011; Prinz et al., 2011; Prinz and Priller, 2014).

### Nuclear receptor agonist treatment of microglia increases phagocytosis

We and others have reported that nuclear receptors act broadly to ameliorate AD-related deficits in murine models of the disease (Skerrett et al., 2014). The agonists lower amyloid burden, and the reduction in amyloid levels was correlated with increased numbers of A $\beta$ -containing myeloid cells in mice treated with nuclear receptor agonists. This led us to test whether Type II nuclear receptor agonists could directly stimulate phagocytosis. Phagocytosis is the process by which microglia and other macrophages take up large (defined as >1  $\mu\text{m}$ ) particles (Aderem and Underhill, 1999). We treated the immortalized N9 microglial cell line with vehicle, the RXR agonist bexarotene (bex), the PPAR $\gamma$  agonist pioglitazone (pio), the LXR agonist GW3965, or the PPAR $\delta$  agonist GW0742 for 18 h before testing the cells' phagocytic capacity with polystyrene beads (Fig. 1B). The nuclear receptor agonists increased basal levels of phagocytosis (Fig. 1B). As has been previously reported, incubation with fibrillar amyloid (Paresce et al., 1996; Bamberger et al., 2003; Reed-Geaghan et al., 2009; Stewart et al., 2010) also stimulated N9 cell phagocytosis (Fig. 1A,C). Furthermore, pretreatment of N9 microglia with nuclear receptor agonists increased their phagocytic response to fibrillar amyloid (Fig. 1A,C) and the TAM receptor ligand Gas6, an adaptor molecule that recognizes apoptotic cells (Fig. 1D). These data demonstrate that nuclear receptor agonists directly enhance the phagocytic activity of N9 cells, as does activation of the TAM receptors.

### Nuclear receptor agonist treatment increases expression of MerTK and Axl *in vitro*

The receptor tyrosine kinases MerTK and Axl are direct genetic targets of nuclear receptor activation in macrophages (A-Gonzalez et al., 2009; Röszer et al., 2011; Daniel et al., 2014). Given that we observed an elevated phagocytic response to the TAM ligand Gas6, we next sought to determine whether this



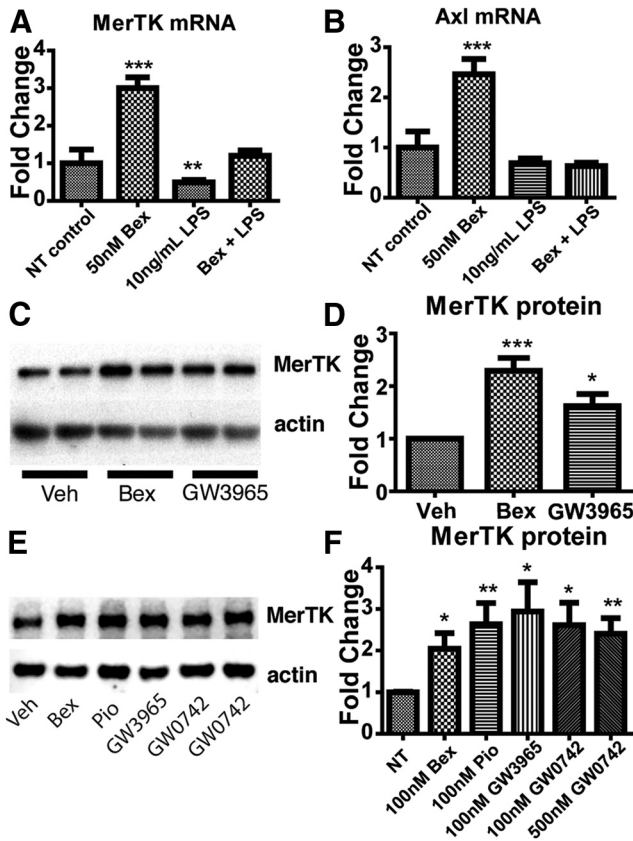
**Figure 1.** Bexarotene increases microglial phagocytosis *in vitro*. N9 microglia were incubated with 1  $\mu\text{m}$  polystyrene beads, and their uptake was measured with flow cytometry (A). N9 cells were treated with or without nuclear receptor agonists (50 nM Bex, RXR agonist; 100 nM pio, PPAR $\gamma$  agonist; 100 nM GW3965, LXR agonist; GW0742, PPAR $\delta$  agonist) for 18 h before incubation with 1  $\mu\text{m}$  polystyrene beads, and the fold change in phagocytic index was measured (B). fA $\beta$  (2.5  $\mu\text{M}$ ) was added 30 min before bead uptake (C). Gas6 (40 nM) was added 30 min before bead uptake (D). Data represent three separate experiments; 10,000 cells were analyzed via flow cytometry. \**p* < 0.05. \*\**p* < 0.01. \*\*\**p* < 0.001.

effect was directly mediated through their induction by nuclear receptor agonists.

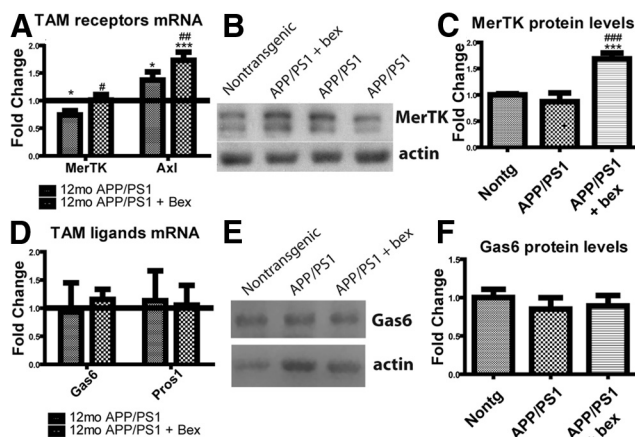
We treated cultured N9 microglia with the RXR agonist bexarotene for 18 h and found that treated microglia expressed higher levels of MerTK and Axl mRNA (Fig. 2A,B) than naive microglia. Treating microglia with the proinflammatory polarizing agent LPS reduced MerTK mRNA levels, and pretreatment with bexarotene was able to prevent this reduction (Fig. 2A). Interestingly, LPS treatment did not affect Axl mRNA levels, although LPS treatment reversed nuclear receptor mediated increases in Axl mRNA expression (Fig. 2B). We next examined MerTK protein levels in microglia treated with the RXR agonist bexarotene or the LXR agonist GW3965. Microglia treated with either bexarotene or GW3965 showed increased levels of MerTK protein expression (Fig. 2C,D). Furthermore, N9 microglia treated with bexarotene, GW3965, the PPAR $\gamma$  agonist pioglitazone, or the PPAR $\delta$  agonist GW0742 also showed increased levels of MerTK protein (Fig. 2E,F). These data provide direct evidence that microglial MerTK expression is regulated by nuclear receptors and this response is sensitive to the inflammatory status of the cells and their environment.

### Bexarotene treatment increases MerTK and Axl expression in two different models of AD

We next investigated the ability of bexarotene to regulate MerTK levels *in vivo*. Interestingly, mRNA levels of MerTK are significantly lower in microglia isolated from 12-month-old APP/PS1 $\Delta\text{E9}$  mice than control mice (Fig. 3A). To determine whether bexarotene could regulate MerTK levels *in vivo*, we treated 12-month-old APP/PS1 $\Delta\text{E9}$  mice for 7 d with 100 mg/kg/d bexarotene or vehicle by oral gavage. Indeed, MerTK mRNA levels were returned to nontransgenic levels after 7 d of bexarotene treatment



**Figure 2.** Nuclear receptor agonist treatment increases MerTK and Axl receptor expression in microglia. At 24 h after addition of bexarotene, N9 microglia were incubated in the presence or absence of 10 ng/ml LPS for an additional 18 h before analysis of MerTK mRNA levels (**A**) and Axl mRNA levels (**B**). Protein levels of MerTK in primary microglia were evaluated by Western analysis 18 h after treatment with 50 nM bexarotene or 500 nM GW3965 (**C**, quantified in **D**). Protein levels of MerTK in N9 microglia were probed via Western blot 18 h after treatment with 50 nM bexarotene, 100 nM pioglitazone, 100 nM GW3965, or 100 or 500 nM GW0742 (**E**, quantified in **F**). Data represent three separate experiments. \* $p < 0.05$ . \*\* $p < 0.01$ . \*\*\* $p < 0.001$ .



**Figure 3.** Bexarotene treatment increases MerTK but not Gas6 expression in APP/PS1 $\Delta$ e9 mice. APP/PS1 $\Delta$ e9 mice (12mo) were treated with 100 mg/kg/d bexarotene or vehicle for 7 d by oral gavage before tissue collection. Microglia were isolated using a Percoll gradient, and MerTK and Axl mRNA levels were investigated using qRT-PCR (**A**). Brain homogenate was probed for MerTK protein levels by Western blot (7 or 8 animals/group) (**B**, quantified in **C**). Levels of Gas6 and Protein S mRNA in microglia isolated from 12-month-old APP/PS1 $\Delta$ e9 animals treated with bexarotene or vehicle for 7 d before tissue collection were investigated using qRT-PCR (**D**). Brain homogenate was probed for Gas6 protein levels by Western blot (7 or 8 animals per group) (**E**, quantified in **F**). \*Difference from nontransgenic animal. #Difference from nontreated APP/PS1 $\Delta$ e9. \* $p < 0.05$ . \*\* $p < 0.01$ . \*\*\* $p < 0.001$ . # $p < 0.05$ . ## $p < 0.01$ . ### $p < 0.001$ .

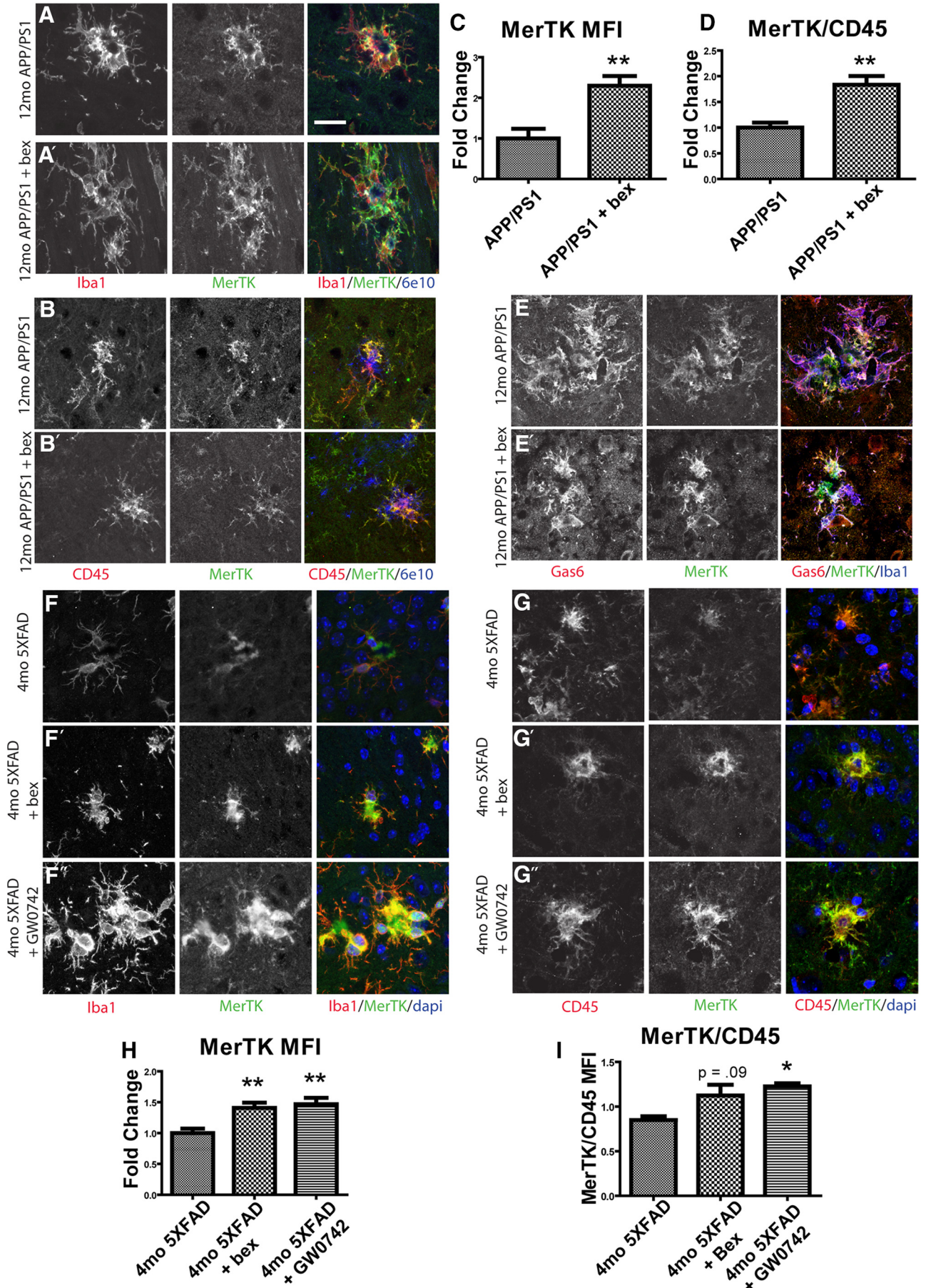
(Fig. 3A). Interestingly, Axl expression was higher in microglia isolated from APP/PS1 $\Delta$ e9 mice than nontransgenic mice and was further increased upon 7 d of bexarotene treatment (Fig. 3A). Furthermore, the levels of MerTK in combined cortical and hippocampal homogenates were measured by Western analysis. Animals treated with bexarotene had significantly higher levels of MerTK than vehicle-treated control animals (Fig. 3B,C).

Because levels of MerTK and Axl were increased by bexarotene treatment of 12-month-old APP/PS1 $\Delta$ e9 mice, we next investigated the effect of genotype and bexarotene treatment on the expression of the two principal TAM receptor ligands, Gas6 and Protein S. Microglia isolated from 12-month-old APP/PS1 $\Delta$ e9 contain similar amounts of Protein S and Gas6 mRNA, regardless of bexarotene treatment (Fig. 3D). However, microglia are not the sole source of TAM ligands within the CNS. We investigated the levels of Gas6 in combined cortical and hippocampal homogenates by Western blot (Fig. 3E, quantified in Fig. 3F) and found no significant differences based on genotype or drug treatment.

In nontransgenic mice, MerTK expression was below the level of detection by immunohistochemistry. However, in both APP/PS1 $\Delta$ e9 and 5XFAD animals, MerTK colocalized with plaque-associated myeloid cells (Fig. 4A,B,E–G). Additionally, MerTK and Gas6 were coexpressed on plaque-associated macrophages in 12-month-old APP/PS1 $\Delta$ e9 animals (Fig. 4E). Significantly, MerTK expression increased in Iba1<sup>+</sup> cells after 7 d of bexarotene treatment (Fig. 4A). Similarly, in plaque-associated cells, MerTK expression was also higher in 4-month-old 5XFAD mice treated with bexarotene or the PPAR $\delta$  agonist GW0742 for 14 d compared with vehicle treated animals (Fig. 4F). These results demonstrate that nuclear receptors increased MerTK expression surrounding plaques in AD mouse models.

To further determine the cell-type specific expression of MerTK, we assessed levels of the marker CD45 on MerTK<sup>+</sup>, plaque-associated cells. CD45 is a canonical marker of myeloid cells (Sedgwick et al., 1991). MerTK colocalized with CD45-expressing cells surrounding amyloid deposits in both 5XFAD and APP/PS1 $\Delta$ e9 animals (Fig. 4B,G). Moreover, those MerTK<sup>+</sup>/CD45<sup>+</sup> cells surrounding plaques were more abundant in 12-month-old APP/PS1 $\Delta$ e9 animals treated with bexarotene (Fig. 4B, quantified in Fig. 4D). Similarly, 4-month-old 5XFAD animals also showed increased ratios of MerTK/CD45 mean fluorescence intensity in response to 14 d of treatment with bexarotene or GW0742 (Fig. 4G, quantified in Fig. 4I). These results further characterize the MerTK<sup>+</sup> plaque-associated cell population that is increased by nuclear receptor agonist treatment.

Because Axl has been shown to have complementary roles to MerTK in mediating phagocytosis (Lemke, 2013), we next investigated whether nuclear receptor treatment would affect Axl expression similarly to MerTK expression *in vivo*. Axl expression was not detected in the nontransgenic brain. However, Axl was abundantly expressed in cells surrounding plaques in both 12-month-old APP/PS1 $\Delta$ e9 and 4-month-old 5XFAD animals (Fig. 5A,C). MerTK and Axl are expressed by the same Iba1<sup>+</sup> cells in APP/PS1 $\Delta$ e9 animals (Fig. 5A). Axl expression was increased in 4-month-old 5XFAD mice treated with bexarotene or PPAR $\delta$  agonist GW0742 for 14 d compared with vehicle-treated animals (Fig. 5D). Similar to what we observed with the expression pattern of MerTK, Axl also colocalized with CD45 expressing cells in both 5XFAD and APP/PS1 $\Delta$ e9 animals (Fig. 5B,E). Upon quantification, 4-month-old 5XFAD animals demonstrate higher Axl/CD45 mean fluorescence intensity in response to 14 d of treatment with either bexarotene or PPAR $\delta$  agonist GW0742 (Fig. 5F). Together, these data support the hypothesis that



MerTK and Axl are similarly regulated by nuclear receptors, and that they're expressed on a common population of Iba1<sup>+</sup>, CD45<sup>+</sup> cells surrounding A $\beta$  deposits.

### TREM2 and MerTK are coexpressed in plaque associated macrophages

TREM2 is expressed exclusively by microglia in the brain, but also by peripheral monocytes and macrophages (Colonna, 2003). We found that the same population of cells in the brain, which are MerTK<sup>+</sup>, CD45<sup>+</sup> also expressed TREM2. Indeed, MerTK and TREM2 colocalize in cells surrounding plaques in both APP/PS1 $\Delta$ e9 (Fig. 6A) and 5XFAD (Fig. 6B) mice.

Jay et al. (2015) have reported that plaque-associated macrophages are derived from blood-borne monocytes that infiltrate the brain and are identified by their high levels of CD45 expression. CD45 is expressed by myeloid cells (Sedgwick et al., 1991), and its expression levels have traditionally been used to discriminate between microglia (CD45<sup>lo</sup>) and peripherally derived myeloid cells (CD45<sup>hi</sup>) (El Khoury et al., 2007; Hickman et al., 2013; Butovsky et al., 2014). Because MerTK and TREM2 colocalized with plaque-associated CD45-expressing cells, we were interested in determining whether MerTK was selectively expressed in either microglia or macrophages derived from infiltrating monocytes. FACS analysis revealed that MerTK expression was largely restricted to the CD45<sup>hi</sup> cells in 5XFAD animals (Fig. 7C). We treated 8-month-old 5XFAD mice with bexarotene or vehicle for 7 d before isolating myeloid cells from the brain. We gated the CD11b<sup>+</sup> population into CD45<sup>lo</sup> and CD45<sup>hi</sup> populations (Fig. 7A). Indeed, 35%–40% of CD11b<sup>+</sup>/CD45<sup>hi</sup> cells in 8-month-old 5XFAD mice expressed TREM2, whereas <1% of CD11b<sup>+</sup>/CD45<sup>hi</sup> cells in nontransgenic littermates were TREM2<sup>+</sup> (Fig. 7B), demonstrating that TREM2 expression is restricted to peripherally derived cells infiltrating the CNS. Less than 5% of CD11b<sup>+</sup>/CD45<sup>lo</sup> cells of any genotype or treatment were TREM2<sup>+</sup> (data not shown), indicating that the resident microglia did not express TREM2 at detectable levels regardless of disease status. There was no effect of bexarotene treatment on the percentage of CD45<sup>hi</sup> cells expressing TREM2 (Fig. 7B). However, MerTK was expressed at a higher level in CD11b<sup>+</sup>/CD45<sup>hi</sup> cells in 5XFAD mice, but not in nontransgenic littermates (Fig. 7C), coincident with MerTK/CD45 coexpression reported on plaque associated cells (Fig. 4B, G). Unsurprisingly, MerTK was expressed at a higher level in TREM2<sup>+</sup> cells as well (Fig. 7D, E), indicating that peripherally derived CD45<sup>hi</sup>, TREM2<sup>+</sup> macrophages also express detectable levels of MerTK. Bexarotene treatment increased both the total MerTK levels, as well as the MerTK levels in TREM2<sup>+</sup> cells (Fig. 7E), indicating RXR agonist treatment increased MerTK levels of both CD45<sup>lo</sup> and CD45<sup>hi</sup> cells in the brains of 5XFAD mice.

←

**Figure 4.** MerTK is expressed by microglia and macrophages in AD model mice and is increased by NR agonist treatment. APP/PS1 $\Delta$ e9 mice (12mo) were treated with vehicle (A, B, E) or the RXR agonist bexarotene (A', B', E') for 7 d by oral gavage. Sections were stained for MerTK (green), Iba1 (red), and amyloid (6e10, blue) (A). MerTK mean fluorescence intensity was quantified (C). MerTK (green) colocalizes with CD45 (red), and the ratio of MerTK/CD45 mean fluorescence intensity was quantified (B, D). MerTK (green) colocalizes with Gas6 (red) and Iba1 (blue) (E). 5XFAD mice (4mo) were treated with vehicle (F, G), bexarotene (F', G'), or the PPAR $\delta$  agonist GW0742 (F'', G'') for 14 d. Sections were stained for MerTK (green) and Iba1 (red), and MerTK mean fluorescence intensity was quantified (F, H). MerTK (green) colocalizes with CD45 (red), and the ratio of MerTK/CD45 mean fluorescence intensity was quantified (G, I). Scale bar, 20  $\mu$ m. Four to six animals were analyzed per group. \* $p$  < 0.05. \*\* $p$  < 0.01.

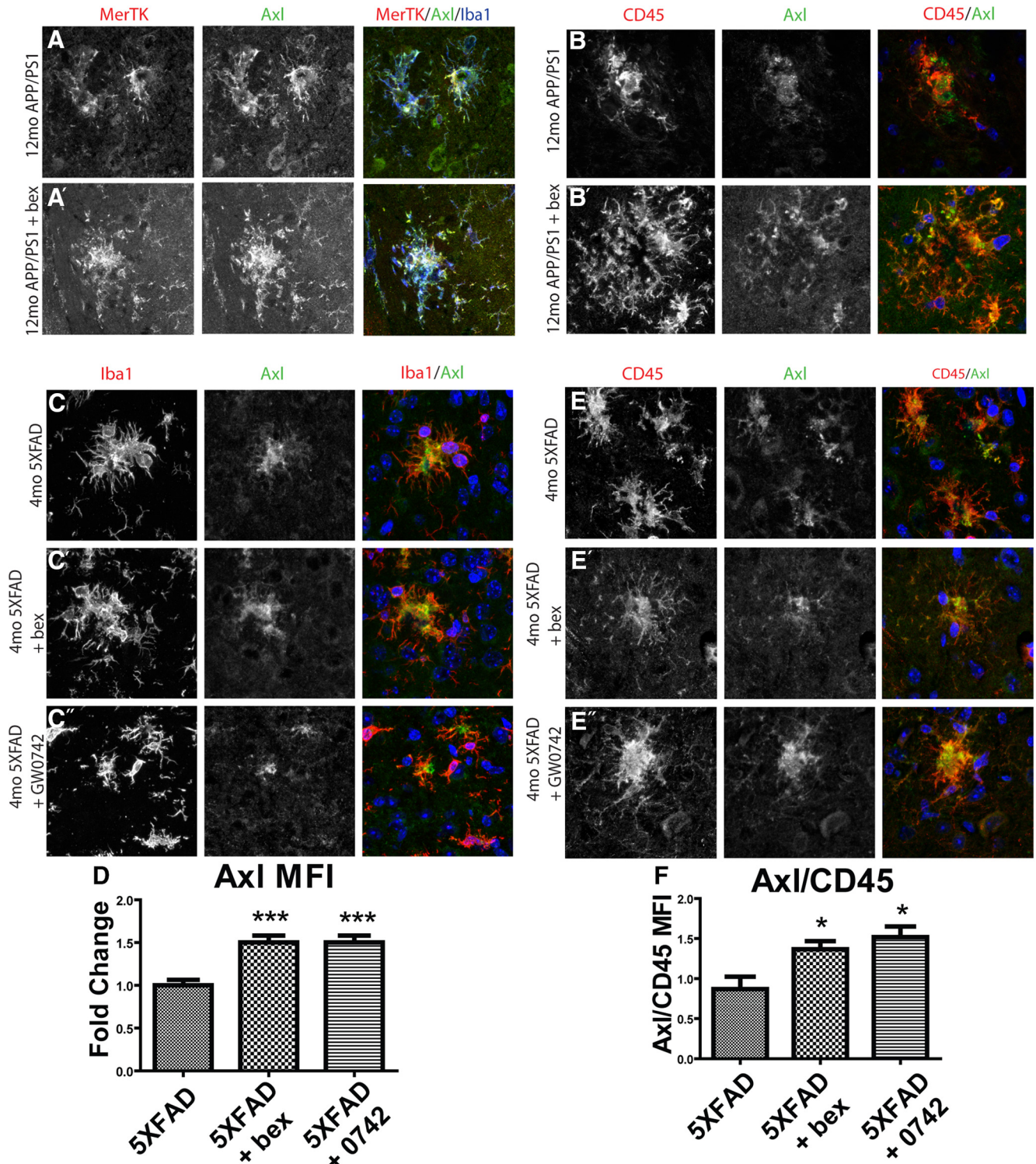
### RXR activation reduces plaque burden in 12 month APP/PS1 $\Delta$ e9 mice and restores phagocytosis in an *ex vivo* phagocytosis assay

We have demonstrated that nuclear receptor treatment both *in vitro* and *in vivo* increased levels of MerTK and Axl and enhanced phagocytosis. We wished to determine whether induction of TAM receptors was necessary for nuclear receptor-mediated increases in phagocytosis. To investigate this potential mechanism, we treated mice with bexarotene and assessed whether modulating signaling through MerTK would alter the ability of bexarotene treatment to enhance phagocytosis in an *ex vivo* phagocytosis assay.

First, we validated that bexarotene treatment induced plaque clearance and enhanced phagocytosis. RXR agonist treatment has been shown to improve behavior and decrease both soluble and fibrillar forms of A $\beta$  in several mouse models of AD (Jiang et al., 2008; Cramer et al., 2012), although others have failed to observe these effects (Tesseur and De Strooper, 2013). We treated 12-month-old APP/PS1 $\Delta$ e9 animals with bexarotene by oral gavage for 7 d. We found acute bexarotene treatment reduced the number of plaques in both cortex and hippocampus by 20% and 30%, respectively, as measured by the number of thioflavin S-positive plaques (Fig. 8A–D). Additionally, insoluble levels of A $\beta$  in combined cortical and hippocampal homogenates were reduced as shown by ELISA (Fig. 8E, F). These data demonstrate that RXR activation reduced plaque burden in aged APP/PS1 $\Delta$ e9 mice over a period of 1 week. The diminished number of plaques was correlated with the appearance of Iba1<sup>+</sup> microglia containing intracellular amyloid in bexarotene-, but not vehicle-treated, mice (Fig. 8G).

Phagocytosis is suppressed by the sustained presence of elevated amyloid levels *in vivo* (Krabbe et al., 2013). Because bexarotene treatment decreased plaque levels and increased macrophage levels of phagocytic receptors MerTK and Axl in a mouse model of AD, we were interested in investigating bexarotene's ability to increase the phagocytic competence of myeloid cells. Cultured microglia are competent phagocytes and readily take up fA $\beta$  introduced into the media (Koenigsknecht and Landreth, 2004). However, these cells are unable to efficiently take up and degrade amyloid in the AD brain, as evidenced by the progressive appearance of insoluble fA $\beta$  plaques throughout the course of the disease. For these reasons, we were interested in studying microglial phagocytosis in a system as close to *in situ* as possible.

We used an *ex vivo* slice phagocytosis assay (Gyoneva and Traynelis, 2013; Krabbe et al., 2013) to determine whether nuclear receptor activation could restore phagocytic competence to microglia residing in the AD brain. We treated 12-month-old APP/PS1 $\Delta$ e9 animals with bexarotene or vehicle for 5 d before preparing brain slices and performing a phagocytosis assay. Indeed, 12-month-old APP/PS1 $\Delta$ e9 animals show reduced phagocytosis compared with nontransgenic littermates (Fig. 9A, B), as previously reported in APPPS1 and APP23 mice (Krabbe et al., 2013). Remarkably, 5 d of oral bexarotene treatment restored phagocytic competence to nontransgenic levels (Fig. 9C). Furthermore, incubation with MerTK and Axl ligand Gas6 restored APP/PS1 $\Delta$ e9 microglial phagocytosis to nontransgenic levels (Fig. 9C). Most strikingly, incubation with a function-blocking anti-MerTK antibody reduced phagocytosis in both nontransgenic slices as well as those from APP/PS1 $\Delta$ e9 animals, which had been treated with bexarotene for 5 d before the assay (Fig. 9C). These data demonstrate that the increase in microglial phagocytosis seen upon bexarotene treatment is reliant on MerTK expression and function.

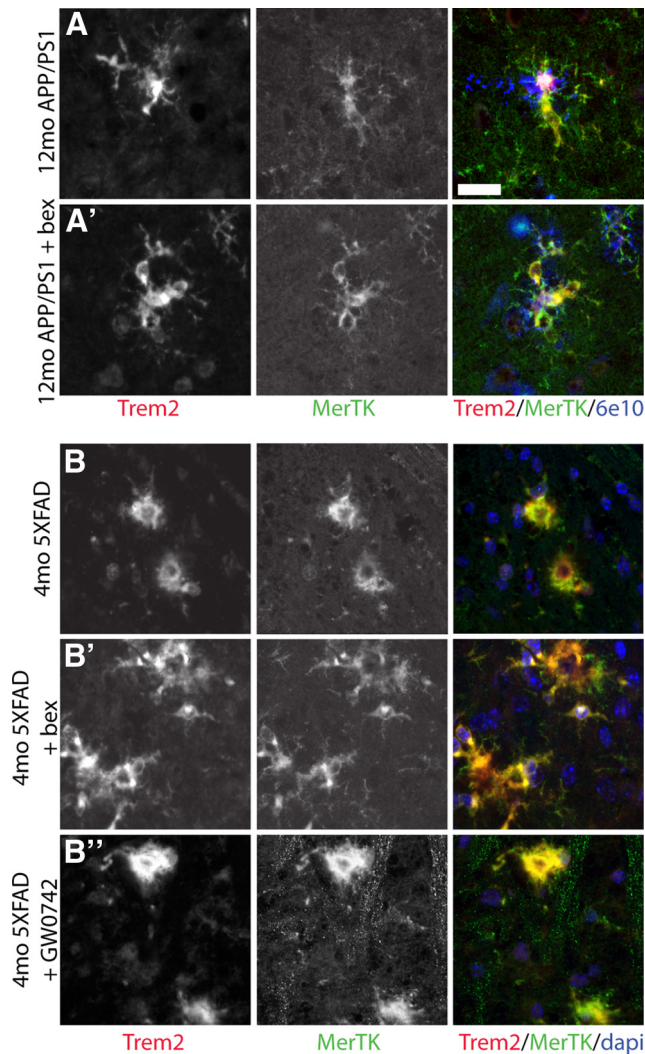


**Figure 5.** Axl is coexpressed with MerTK by microglia and macrophages in AD model mice and is increased with nuclear receptor agonist treatment. APP/PS1Δe9 mice (12mo) were treated with vehicle (*A, B*) or the RXR agonist bexarotene (*A', B'*) for 7 d by oral gavage. Sections were stained for Axl (green), MerTK (red), and Iba1 (blue) (*A*). 5XFAD mice (4mo) were treated with vehicle (*C, E*), bexarotene (*C', E'*), or PPARδ agonist GW0742 (*C, D', E'*) for 14 d before sectioning and staining. Sections were stained for Axl (green) and Iba1 (red), and Axl mean fluorescence intensity was quantified (*D*). Axl (green) colocalizes with CD45 (red) and the ratio of Axl/CD45 mean fluorescence intensity was quantified (*F*). Scale bar, 20 μm. Four to six animals were analyzed per group. \**p* < 0.05. \*\*\**p* < 0.001.

**Discussion**

Understanding of the complex biology of microglia has been hampered by the lack of a conceptual framework through which to interpret how myeloid cell phenotype is regulated in the normal and diseased brain. The triggers of inflammation, such as

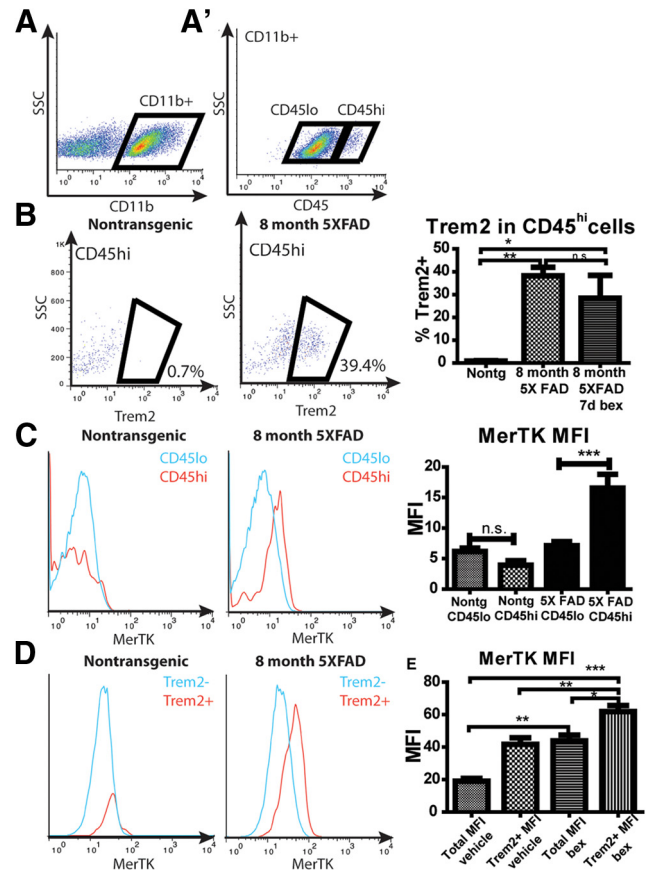
elevated soluble and deposited amyloid levels in the AD brain, are well studied (Akiyama et al., 2000). However, the chronic adaptations of myeloid cells to the diseased brain environment are not well understood. A central role for inflammation in the pathogenesis of Alzheimer’s disease and other neurodegenerative dis-



**Figure 6.** MerTK and TREM2 are expressed by plaque-associated macrophages. MerTK (green) colocalizes with TREM2 (red) and surround  $6e10^+$  plaques (blue) in 12-month-old APP/PS1 $\Delta e9$  mice treated with vehicle (**A**) or bexarotene (**A'**) for 7 d. MerTK (green) also colocalizes with TREM2 (red) in 4-month-old 5XFAD mice treated with vehicle (**B**), bexarotene (**B'**), or GW0742 (**B''**). Scale bar, 20  $\mu$ m.

eases has been buttressed by a succession of studies identifying genetically linked risk factors that regulate or affect the activity of immune cells in the brain (Hollingworth et al., 2011; Naj et al., 2011). Indeed, the cell biological evidence, embodied within a large literature (Akiyama et al., 2000), and in directed studies focused on newly identified risk genes, such as TREM2, CD33, CD300a, and others, is largely consistent with the actions of immune responses in the brain that confer elevated disease risk (Jones et al., 2010; Guerreiro et al., 2013; Jonsson et al., 2013).

One of the unexplained aspects of AD pathogenesis is why microglia fail to effectively phagocytose fibrillar amyloid deposits, reflecting the disengagement of the phagocytic machinery directed toward this target. These cells express an array of receptors that act in an ensemble to recognize fibrillar forms of amyloid, including Scara1, CD47,  $\alpha 6\beta 1$  integrin, TLRs 2/4/6, and their coreceptors CD36 and CD14 (Paresce et al., 1996; Bamberger et al., 2003; Reed-Geaghan et al., 2009; Stewart et al., 2010). Importantly, microglia in the AD brain are phagocytically competent, as opsonization of amyloid with antibodies (Bard et al., 2000; Demattos et al., 2012) or



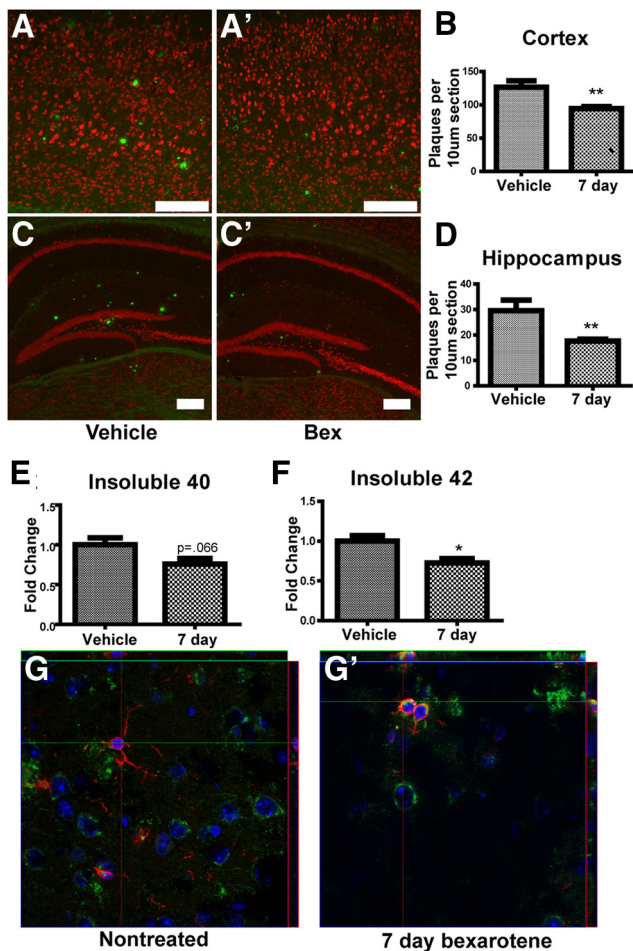
**Figure 7.** MerTK and TREM2 are expressed by peripherally derived myeloid cells. Microglia were isolated from 8-month-old 5XFAD mice (4–6 per group) and gated on CD11b $^+$ /CD45 $^{lo}$  and CD11b $^+$ /CD45 $^{hi}$  cell populations (**A**, **A'**). **B**, TREM2 expression in the CD11b $^+$ /CD45 $^{hi}$  cells was quantitated. **C**, Mean fluorescence intensity of MerTK was measured in CD11b $^+$ /CD45 $^{lo}$  and CD11b $^+$ /CD45 $^{hi}$  populations in both transgenic and nontransgenic mice. Mean fluorescence intensity of MerTK was measured in the TREM2 expressing population (**D**, quantified in **E**). \* $p < 0.05$ . \*\* $p < 0.01$ . \*\*\* $p < 0.001$ . n.s., Not significant.

complement (Fu et al., 2012) can effectively stimulate phagocytosis and plaque clearance. The presence of amyloid acts to suppress macrophage phagocytic function (Krabbe et al., 2013), whereas the interaction of macrophages with plaques elicits the secretion of an array of inflammatory cytokines, which directly suppress phagocytosis (Koenigsnecht-Talboo and Landreth, 2005).

The present study describes a mechanism whereby macrophages in the AD brain are induced to phagocytose and clear amyloid deposits through the action of nuclear receptors, which regulate the expression of arrays of functionally allied genes to license this response. Phagocytes use a diverse cohort of receptors that recognize and engulf cellular fragments and apoptotic cells whose membrane lipids serve as direct ligands of the nuclear receptors. However, until now, it has not been clear exactly how various nuclear receptors mediate enhanced phagocytosis. Here, we show that the enhanced phagocytic capacity is subserved by the induction of the phagocytic receptors Axl and MerTK, as well as a cohort of other genes supporting this function (Daniel et al., 2014).

There is an extensive literature on the salutary actions of nuclear receptors in murine models of AD and other CNS disorders, where they act to promote the conversion of microglia from pro-inflammatory to anti-inflammatory phenotypes (Skerrett et al., 2014). Studies using PPAR $\gamma$ , PPAR $\delta$ , LXR, and RXR agonists have consistently reported amelioration of AD-related phenotypes

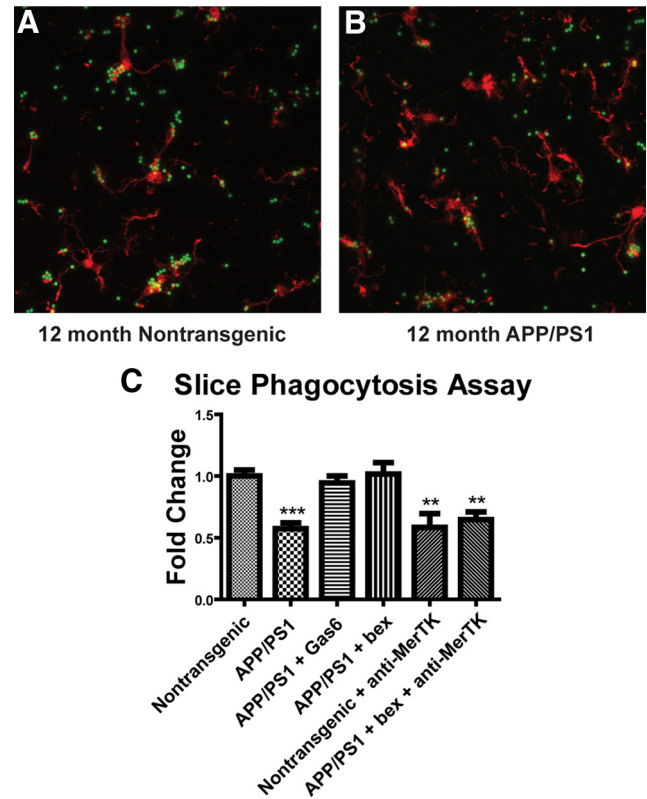




**Figure 8.** Bexarotene treatment decreases plaque burden in APP/PS1Δe9 animals. APP/PS1Δe9 mice (12 months) were treated with vehicle (**A**, **C**) or 100 mg/kg/d bexarotene (**A'**, **C'**) for 7 d by oral gavage before staining with Thioflavin S. Plaques in the cortex (**B**) and hippocampus (**D**) were counted from 5 animals. Scale bar, 200 μm. ELISAs of Aβ40 and Aβ42 were run on formic acid extractions of cortical homogenates from 7 or 8 animals per group to determine insoluble Aβ40 and Aβ42 levels (**E**, **F**). Confocal images show Aβ (6e10, green) inside microglia (Iba1, red) (**G**). \* $p < 0.05$ . \*\* $p < 0.01$ .

and behavioral impairments. PPAR $\gamma$  and RXR have been shown to increase phagocytosis of amyloid in a CD36-dependent mechanism (Yamanaka et al., 2012), as well as control C1 and Fc $\gamma$ R transcript levels on peripheral macrophages (Röszer et al., 2011). Plaque reduction was observed in the majority of these studies but was the most variable outcome and is influenced by age and aggressiveness of the disease models. The nuclear receptor agonists in these studies were typically delivered chronically through the diet over a period of months. We demonstrated that acute (3–9 d) treatment with PPAR $\gamma$  (Mandrekar-Colucci et al., 2012) or RXR agonists (Cramer et al., 2012) was sufficient to induce plaque loss, but this effect was highly sensitive to drug formulation and pharmacodynamics (Landreth et al., 2013; Tesseur and De Strooper, 2013; Chen et al., 2014). The present work identifies a more complex biology associated with the phagocytic clearance of amyloid deposits from the brain because of the identification of the peripheral origins of the plaque-associated macrophages and their responses to nuclear receptor agonists within the brain.

We report here and in related work (Jay et al., 2015) that plaque-associated macrophages express markers consistent with myeloid cells derived from peripheral monocytes. Blood-borne 'inflammatory' monocytes are CD45<sup>hi</sup>, whereas microglia are



**Figure 9.** Bexarotene treatment increases phagocytosis in 12-month-old APP/PS1Δe9 mice in a MerTK-dependent manner. The 12-month-old APP/PS1 mice were treated with 100 mg/kg/d bexarotene or vehicle for 5 d by oral gavage before *ex vivo* slice culture phagocytosis assay. Slices were taken from nontransgenic (**A**) or APP/PS1Δe9 (**B**) animals treated with 100 mg/kg/d bexarotene or vehicle and incubated with Gas6 or function blocking anti-MerTK antibody and 2 μm polystyrene beads to determine their phagocytic index (**C**). Data represent three separate experiments. \*\* $p < 0.01$ . \*\*\* $p < 0.001$ .

CD45<sup>lo</sup>. CD45 expression has historically been used to discriminate between these cells, which arise from entirely distinct lineages and more recently has allowed the delineation of transcriptional differences between microglia and other myeloid populations (Gautier et al., 2012; Hickman et al., 2013; Butovsky et al., 2014; Yamasaki et al., 2014). The suggestion that monocytes infiltrate the AD brain and have salutary effects on AD pathogenesis is not new (Simard et al., 2006; El Khoury and Luster, 2008), but these studies were confounded by technical considerations (Morganti et al., 2014). However, El Khoury et al. (2007) and El Khoury and Luster (2008) reported that peripheral myeloid cells infiltrated the AD brain in a CCL2-dependent manner, and loss of CCR2 was associated with fewer parenchymal macrophages, increased plaque burden, and increased levels of Aβ peptides. Our observation that MerTK<sup>+</sup>/Axl<sup>+</sup> plaque-associated macrophages are also CD45<sup>hi</sup> and TREM2<sup>+</sup> argues strongly that these cells are derived from peripheral monocytes. Importantly, our data show that nuclear receptor agonists increase phagocytic receptor expression on both TREM2<sup>-</sup> and TREM2<sup>+</sup> populations and enhance phagocytosis in these different populations.

The question of whether MerTK<sup>+</sup>/Axl<sup>+</sup> plaque-associated macrophages are derived from resident microglia or peripheral monocytes will be important to explore further as they may have distinct roles in disease and be regulated differently by nuclear receptor agonist treatment. The roles of resident microglia and infiltrating monocytes have only recently begun to be delineated (Yamasaki et al., 2014). New work suggests that the action of

nuclear receptors in circulating myeloid subpopulations may license a different array of genes that subserve the emigration of these cells into tissues to sites of inflammation and their subsequent differentiation (Daniel et al., 2014). Analysis of RXR-regulated gene expression in peripheral monocytes has revealed the coordinate induction of genes associated with extravasation through the endothelium (CD300a, *Ninj1*, *CCL2*), differentiation into macrophages (*MafB*, *IRF8*), and regulation of phagocytosis (*TREM2*, *MerTK*, *Axl*, *CD300a*, *C1q*, *CD36*) (Daniel et al., 2014). Nuclear receptor actions in tissue macrophages have been observed on a cohort of genes that are specifically associated with “alternative activation” and phagocytosis (Odegaard et al., 2007; A-Gonzalez et al., 2009; Chinetti-Gbaguidi et al., 2011; Röszer et al., 2013).

*MerTK* and *Axl* are necessary for the phagocytosis of apoptotic cells but also act broadly to regulate the innate immune response in tissue macrophages (Lemke, 2013). These receptors exhibit important anti-inflammatory actions, and their expression serves as a marker of “alternative activation” (Zizzo et al., 2012). *Axl/MerTK* have recently been shown to be expressed in a mutually exclusive manner on bone marrow-derived macrophages. *MerTK* acts in a homeostatic role and is induced in immunosuppressive environments, whereas *Axl* levels are increased by proinflammatory factors (Zagórska et al., 2014). Both receptors act as negative feedback regulators to reduce inflammation (Rothlin et al., 2007); however, neither has been studied in the context of AD. We found that plaque-associated macrophages express high levels of both *MerTK* and *Axl* in the AD brain, which are further increased through the activation of nuclear receptors. Importantly, our data suggest that *MerTK* activity is necessary for amyloid-stimulated phagocytosis, and it will be important to understand the mechanistic basis of this phenomenon.

*TREM2* has classically been considered a phagocytic receptor, based largely on *in vitro* studies demonstrating its requirement for the phagocytosis of apoptotic neurons and bacteria. However, we demonstrate that *TREM2*<sup>+</sup> cells accumulate on plaques where they exhibit an inflammatory phenotype and are phagocytically ineffective, as evidenced by the progressive increase in plaque burden throughout the course of the disease. Similarly, it is widely held that *TREM2* acts to suppress inflammatory gene expression, based largely on data demonstrating that knockdown or genetic knock-out of *TREM2* resulted in higher levels of proinflammatory cytokines (Takahashi, 2005; Turnbull et al., 2006; Sharif et al., 2014). This view is inconsistent with the observation of inflammatory macrophages associated with plaques, virtually all of which express high levels of *TREM2*. *TREM2* has recently been reported to have analogous proinflammatory actions in murine models of stroke (Sieber et al., 2013), colitis (Correale et al., 2013), and lung inflammation (Sharif et al., 2014). Importantly, Jay et al. (2015) have shown a dramatic decrease in monocyte infiltration in *TREM2*<sup>-/-</sup> AD model mice and an absence of plaque-associated cells, suggesting that *TREM2* has previously unappreciated roles in peripheral monocytes and their subsequent actions within the brain.

This study provides direct evidence that nuclear receptors act to promote the clearance of amyloid deposits from the brain, in part, through their ability to induce the expression of the phagocytic receptors *Axl* and *MerTK* on plaque-associated macrophages. The recognition that these latter cells are likely derived from circulating monocytes and express *TREM2* force a substantial reevaluation of both the actions of *TREM2* and the roles of myeloid cells in AD pathogenesis.

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